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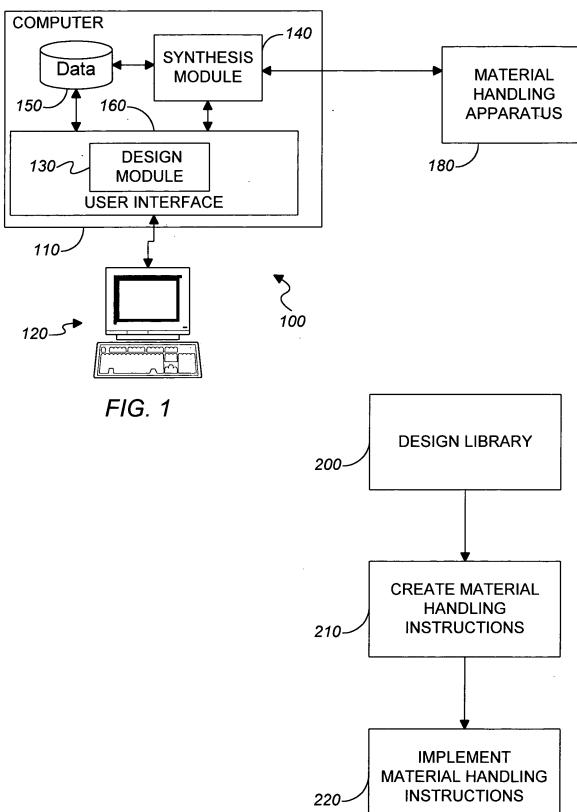


FIG. 2

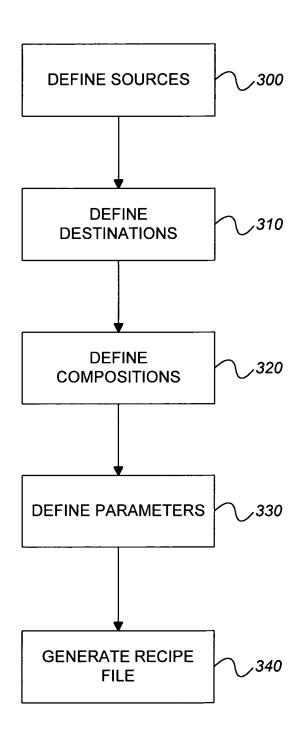
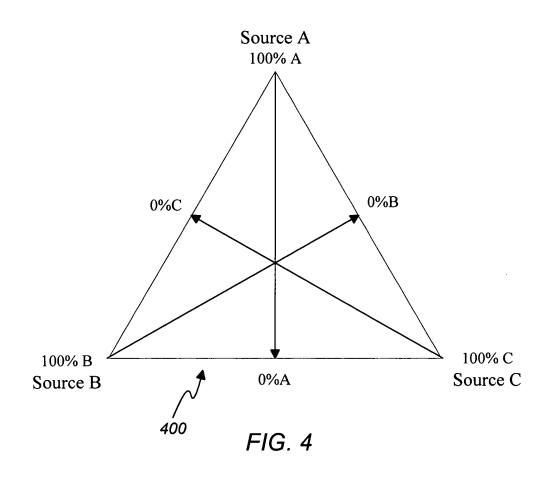


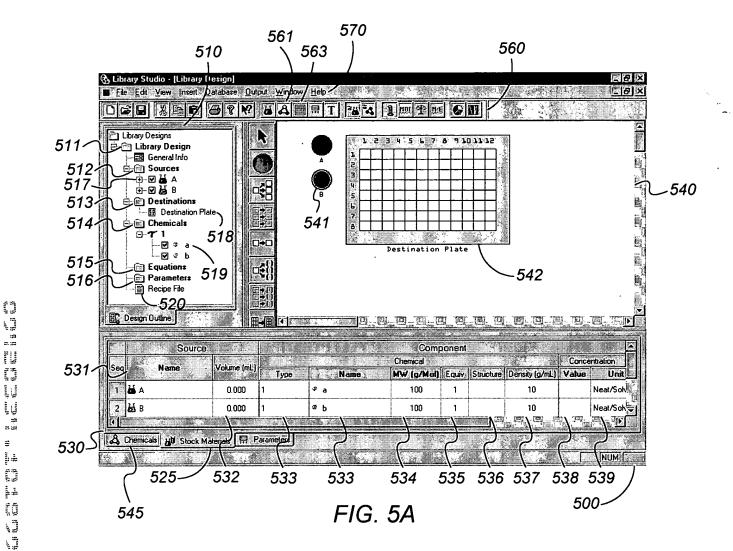
FIG. 3

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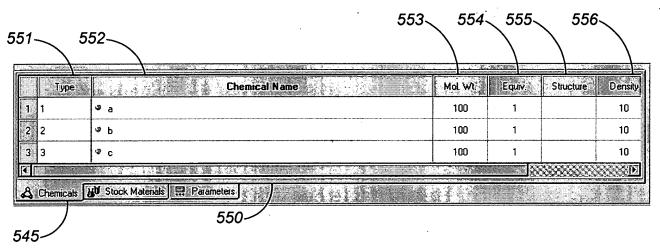


FIG. 5B

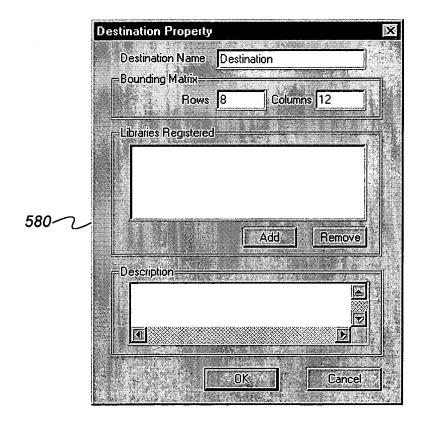


FIG. 5C

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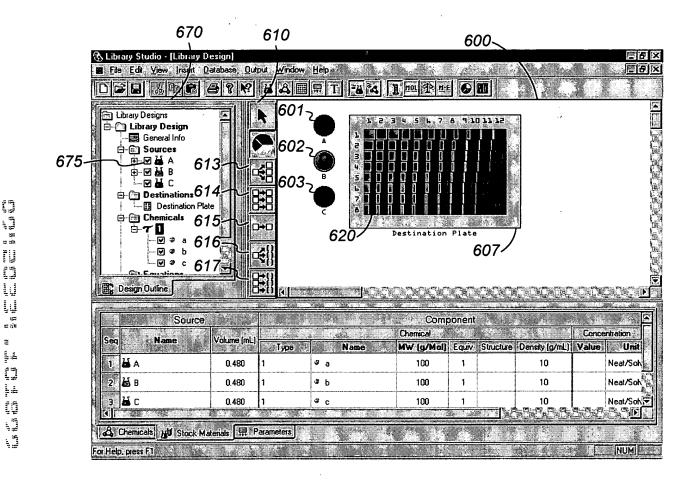


FIG. 6A

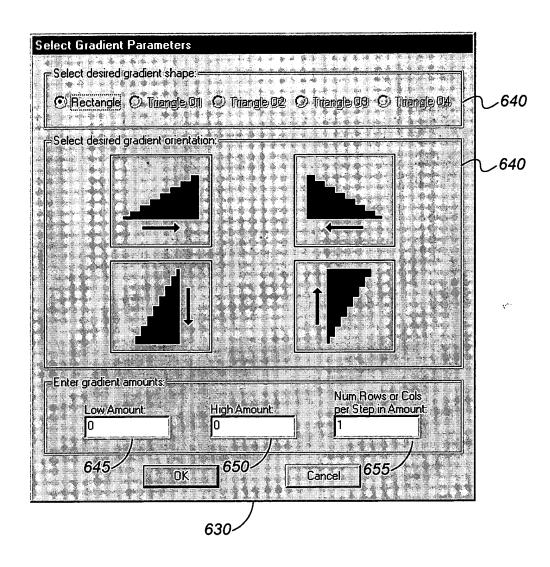


FIG. 6B

pping	Sequence							3 3 8
Seqt	Source	De	kination	Amount	To Recipe	Tag 🧖	Collaboration	ert
1	F (1,1),(1,1) Pla	te (1,1),(7,4)	10.00 to 70.00	☑	Ĭ.		
2	E (1,1),(1,1) Pla	te (1,5),(7,8)	10.00 to 70.00	$oldsymbol{arPsi}$		>>:De	lete
L 3	G (1,1),(1,1) Pla	te (1,9),(7,12)	10.00 to 70.00	v		1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	
4	B (1,1),(1,1) Pla	te (1,1),(2,4)	10.00 to 40.00	V		Modify	
4 5	B (1,1),(1,1) Pla	te (1,5),(2,8)	10.00 to 40.00	2		Replic	ate
- 6	B (1,1),(1,1) Pla	te (1,9),(2,12)	10.00 to 40.00	v			
4 7	C (1,1),(1,1) Pla	te (5,1),(6,4)	10.00 to 40.00	v	8		
-4 8	C (1,1),(1,1) Plai	te (5,5),(6,8)	10.00 to 40.00	9			
- 9	C (1,1),(1,1) Pla	te (5,9),(6,12)	10.00 to 40.00	v	Ž.		
10	B (1,1),(1,1) Pla	te (8,1),(8,2)	70.00 to 70.00	Y			
11	C (1,1),(1,1) Plai	te (8,9),(8,10)	70.00 to 70.00	y	•		
12	H (1,1),(1,1) Pla	te (1,1),(8,12)	500.00 to 500.0	V	į.	Caramanana	
4 13	L[1,1],[1],A) Pla	e (3,1),(4,4)	10.00 to 40.00	7		- 	

FIG. 6C

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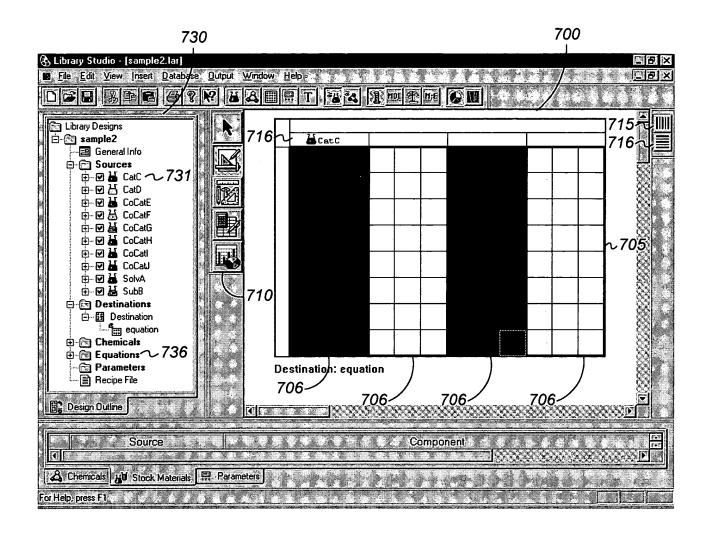


FIG. 7A

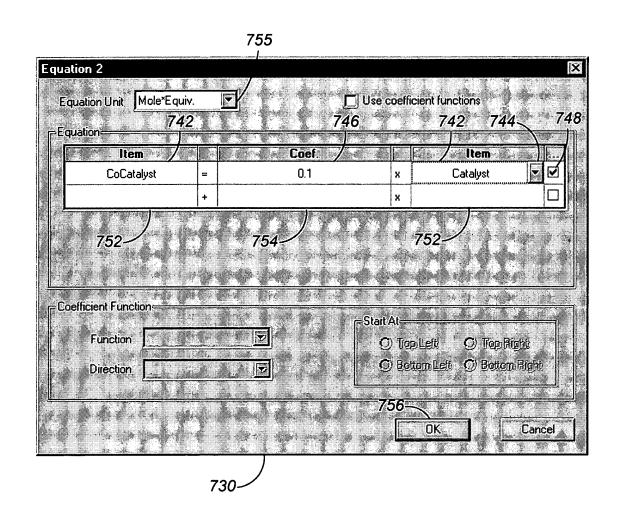


FIG. 7B

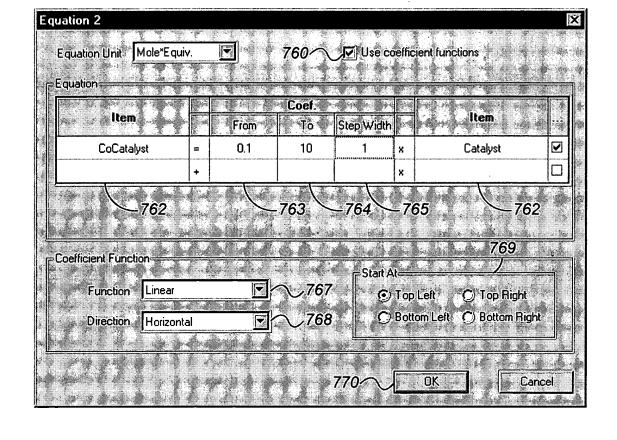
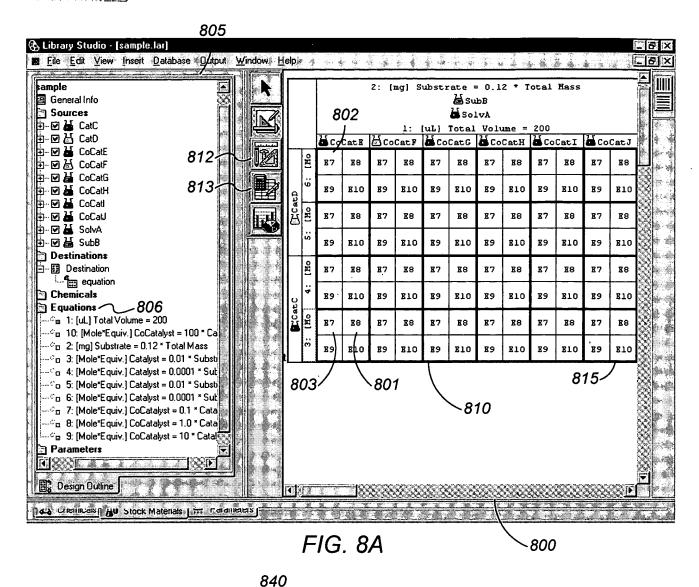


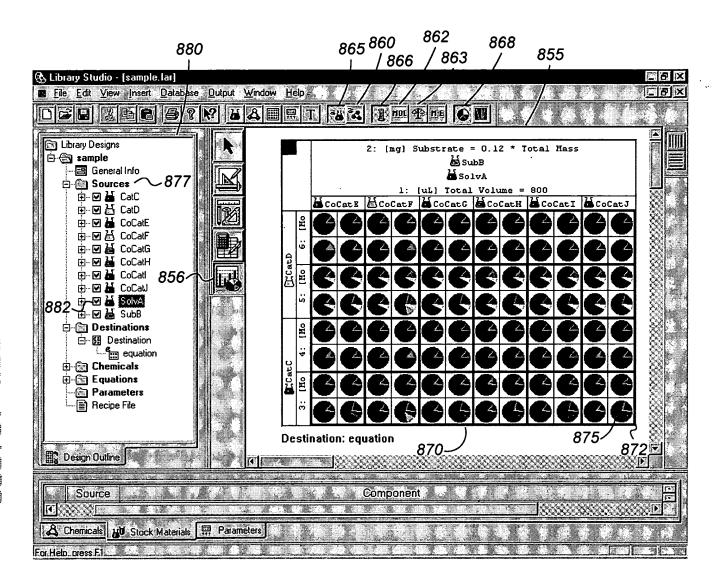
FIG. 7C





¿ Equation Matrixes - Cell (1, 1) atus: Equation solving failed SubB CoCatE CatO Mole Equiv I CoC 0.000000 0.000000 -0.000001 0.000010 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 20.00000 (mg) mg) Substrate = 0. -0.120000 0.880000 -0.120000 -0.120000 -0.120000 0.000000 0.000000 0.000000 0.000000 0.000000 2.72739 (mg) 0.000000 1.000000 0.00027 (mg) (uL) Total Volume 0.000000 0.000000 0.000000 0.000000 1.0000000 1.000000 1.000000 200.000000 Mole Equiv.] Cata 0.000000 -0.000000 0.000010 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.00003 (mg) 0.00055 (mg) 0.000000 0.000000 0.000000 -1.0000000 0.000000 0.000000 0.100000 0.100000 0.100000 0.100000 0.000000 -1.0000000 0.000000 0.000000 0.000000 0.001000 0.000000 0.000000 0.000000 0.000000 2727.39 (uL 0.000000 0.000000 0.000000 0.000000 1.000000 0.000000 0.000000 0.000000 0.001000 0.000000 SolvA -2527.66 (uL 0.000000 0.000000 0.000000 -1.000000 0.000000 0.000000 0.000000 0.010000 0.000000 0.000000 CoCatE 0.00 (ut. 0.000000 0.000000 0.000000 0.000000 0.000000 0.27 (uL 0.000000 -1.000000 0.000000 0.000000 0.002000 OK. 835 830 845 850

FIG. 8B



- 32 S. S. S.

FIG. 8C

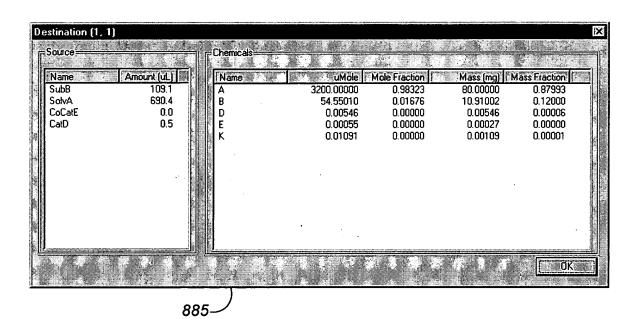


FIG. 8D

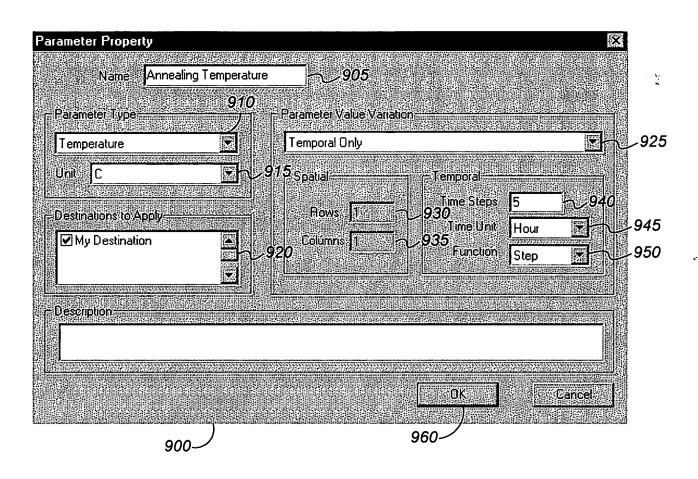


FIG. 9A

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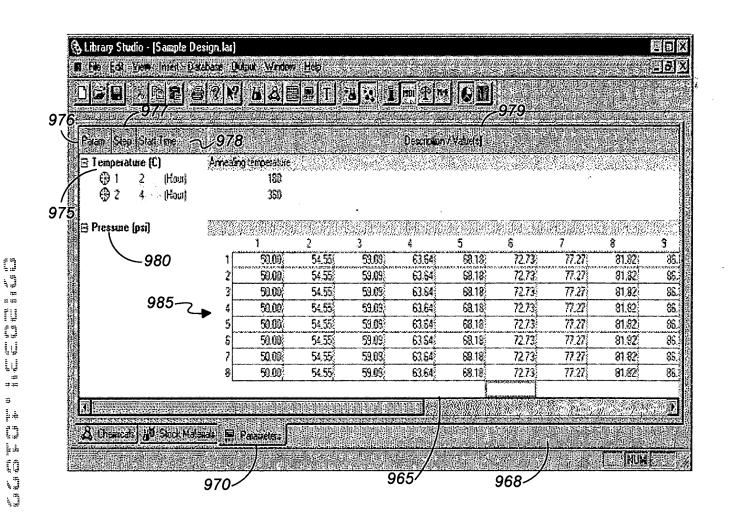


FIG. 9B

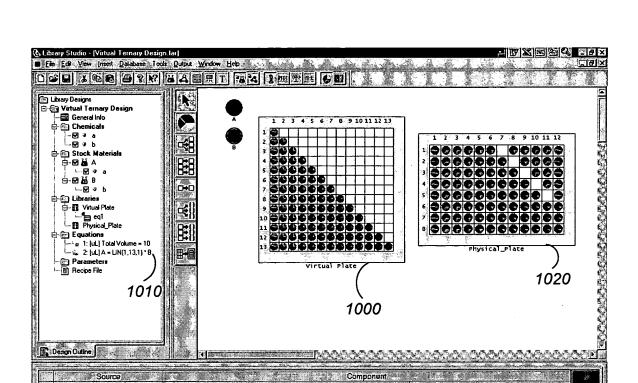


FIG. 10

1

Type MW (g/Mail) Equiv Structure Density (g/mL) Velue Unit Assumption Core (g/mL)

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